SEQUENCE LISTING

<110> Voorberg, Johannes

<120> Method For Diagnosis and Treatment of Haemophilia A Patients With An Inhibitor

<130> Sequence Nos 1-55 for 294-86 PCT/US/RCEII

<140> 09/674,752
<141> 2000-12-29

<150> PCT/NL99/00285
<151> 1999-05-07

<150> EP 98201543.0
<151> 1998-05-08

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<170> PatentIn version 3.3

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35 40 45

Gly Gly Ile Ile Pro Ile Phe Gly Ser Thr Lys Tyr Ala Gln Lys Phe
50 55 60

Gln Gly Arg Val Thr Met Thr Ala Asp Gly Ser Thr Ser Thr Ala Tyr
65 70 75 80

Met Glu Leu Asn Ser Leu Arg Ser Glu Asp Thr Ala Ile Tyr Tyr Cys
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Ala Ile Ser Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
35 40 45

Gly Gly Ile Ile Pro Ile Phe Gly Thr Ala Asn Tyr Ala Gln Lys Phe
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Gln Gly Arg Val Thr Ile Thr Ala Asp Glu Ser Thr Ser Thr Ala Tyr
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Ala Arg

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35 40 45

Gly Trp Ile Ser Ile Tyr Ser Gly Asn Thr Asp Tyr Ala Gln Lys Phe
50 55 60

Gln Gly Arg Val Thr Met Thr Thr Asp Thr Ser Arg Arg Thr Ala Tyr
65 70 75 80

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85 90 95

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35 40 45

Gly Trp Ile Ser Ala Tyr Asn Gly Asn Thr Asn Tyr Ala Gln Lys Leu
50 55 60

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65 70 75 80

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85 90 95

Ala Arg

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20 25 30

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35 40 45

Gly Trp Ile Ser Ala Tyr Asn Gly Asn Thr His Tyr Ala Gln Lys Phe
50 55 60

Gln Gly Arg Val Thr Met Thr Thr Asp Thr Ser Arg Arg Thr Ala Tyr
65 70 75 80

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Ala Arg

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Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Pro Phe Thr Ser Tyr
20 25 30

Asp Ile Ser Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
35 40 45

Gly Trp Ile Ser Ile Tyr Ser Gly Asn Thr Asp Tyr Ala Gln Lys Phe
50 55 60

Gln Gly Arg Val Thr Met Thr Thr Asp Thr Ser Arg Arg Thr Ala Tyr
65 70 75 80

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20 25 30

Asp Ile Asn Trp Val Arg Gln Ala Thr Gly Gln Gly Leu Glu Trp Met
35 40 45

Gly Trp Met Asn Pro Asn Ser Gly Asn Thr Gly Tyr Ala Gln Lys Phe
50 55 60

Gln Gly Arg Val Thr Met Thr Arg Asn Thr Ser Ile Ser Thr Ala Tyr
65 70 75 80

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85 90 95

Ala Arg

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20 25 30

Asp Ile Asn Trp Val Arg Gln Ala Thr Gly Gln Gly Leu Glu Trp Met
35 40 45

Gly Trp Met Asn Pro Asn Ser Gly Asn Ala Gly Phe Ala Gln Lys Phe
50 55 60

Lys Gly Arg Leu Thr Leu Thr Arg Asp Thr Ser Thr Ser Thr Ala Tyr
65 70 75 80

Met Glu Leu Arg Asn Leu Glu Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Arg Cys Asp Thr Thr Leu Leu Ile Trp Phe Gly Pro Ala Pro Tyr
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Asn Asp Ser Trp Gly Gln Gly Thr Leu Val
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20 25 30

Ala Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Gly Ile Ser Trp Asn Ser Gly Ser Ile Gly Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Ser Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Leu Tyr Tyr Cys
85 90 95

Ala Lys Asp

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20 25 30

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35 40 45

Ser Gly Val Thr Trp Ser Gly Thr Thr Ile Gly Phe Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Ser Leu Tyr
65 70 75 80

Leu Tyr Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Leu Tyr Tyr Cys
85 90 95

Ala Leu Pro Tyr Ile Asn Ser Ser Asn Tyr Arg Arg Gly Val Ala Ala
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
20 25 30

Gly Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ala Val Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala Asp Ser Val

50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Lys

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20 25 30

Gly Met His Trp Val Arg Gln Ala Pro Gly Ala Gly Leu Glu Trp Val
35 40 45

Ala Val Ile Ser Tyr Asp Gly Asn Asp Lys Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Ala Ile Ser Arg Asp Asn Ala Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Thr Ile Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Lys Asp Leu Ile Glu Ser Asn Ile Ala Glu Ala Leu Trp Gly Gln
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Gly Thr Leu Val Thr Val Ser Ser
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<213> Homo sapiens

<400> 37

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            20                  25                  30

Ser Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
        35                  40                  45

Ser Ser Ile Ser Ser Ser Ser Ser Tyr Ile Tyr Tyr Ala Asp Ser Val
    50                  55                  60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Ser Leu Tyr
65                  70                  75                  80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
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            20                  25                  30

Asp Ile His Trp Val Arg Gln Thr Pro Gly Lys Gly Leu Glu Trp Val
        35                  40                  45

Ser Ser Ile Ser Ser Gly Gly Asn Tyr Ile Asp Tyr Ala Asp Ser Val
    50                  55                  60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Asn Asn Val Val Tyr
65                  70                  75                  80
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Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Met Ala Val Tyr Phe Cys
                85                  90                  95

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gcacagaagt ttaagggcag actcaccttg accagggaca cttccacaag cacagcctac    240

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Asp Ile Asn Trp Val Arg Gln Ala Thr Gly Gln Gly Leu Glu Trp Met
        35                  40                  45

Gly Trp Met Asn Pro Asn Ser Gly Asn Ala Gly Phe Ala Gln Lys Phe
    50                  55                  60
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Lys Gly Arg Leu Thr Leu Thr Arg Asp Thr Ser Thr Ser Thr Ala Tyr
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Met Glu Leu Arg Arg Leu Glu Ser Glu Asp Thr Ala Val Tyr Tyr Cys
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Ala Arg Cys Asp Thr Thr Leu Leu Ile Trp Phe Gly Pro Ala Pro Tyr
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<400> 47

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20 25 30

Asp Ile His Trp Val Arg Gln Thr Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Ser Ile Ser Ser Gly Gly Asn Tyr Ile Asp Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Asn Asn Val Val Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Met Ala Val Tyr Phe Cys
85 90 95

Ala Arg Asp Gly Thr Ile Phe Gly Ser Ala Ala Thr Trp Arg Ala Phe

100 105 110

Asp Ile Trp Gly Arg Gly Thr Met Val Thr Val Ser Ser
115 120 125

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20 25 30

Ala Ile Ser Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
35 40 45

Gly Asp Ile Ile Pro Ile Leu Gly Thr Gly Asn Tyr Ala Gln Lys Phe
50 55 60

Gln Gly Arg Val Thr Ile Thr Ala Asp Glu Ser Thr Ser Thr Ala Tyr
65 70 75 80

Met Glu Leu Ser Thr Leu Thr Ser Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Glu Leu Asp Trp Phe Tyr Ile Trp Gly Gln Gly Thr Met Val Thr Val
100 105 110

Ser Ser

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<213> Homo sapiens

<400> 50

Glu Val Gln Leu Leu Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
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Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Ser Tyr
20 25 30

Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
65 70 75 80

Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
85 90 95

Ala Lys

<210> 51
<211> 122
<212> PRT
<213> Homo sapiens

<400> 51

Glu Val Gln Leu Val Glu Ser Gly Gly Asp Leu Val Gln Pro Gly Gly
1 5 10 15

Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser Asn Phe

20 25 30

Ala Met Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val
35 40 45

Ala Ala Ile Gly Gly Arg Ser Gly Thr Thr Phe Tyr Ala Asp Ser Val
50 55 60

Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Val Tyr
65 70 75 80

Leu Glu Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Ile Tyr Tyr Cys
85 90 95

Ala Lys Arg Gly Arg Gly Gly Tyr Lys Tyr Tyr Gly Met Asp Val Trp
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Gly Gln Gly Thr Thr Val Thr Val Ser Ser
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<210> 52
<211> 342
<212> DNA
<213> Homo sapiens

<400> 52
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cctggacaag ggcttgagtg gatgggagac atcatcccta tccttggtac aggaaactac 180

gcacagaagt tccagggcag agtcacgatt accgcggacg agtccacgag cacagcctac 240

atggagctga gcaccctgac atctgaggac acggccgtgt attactgtga acttgactgg 300

ttctatatct ggggccaagg gacaatggtc accgtgtcga gt 342

<210> 53
<211> 342
<212> DNA
<213> Homo sapiens

<400> 53
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ctcgtccgcg gtaatcgtga ctctgccctg gaacttctgt gcgtagtttc ctgtaccaag 180

gatagggatg atgtctccca tccactcaag cccttgtcca ggggcctgtc gcacccagct 240

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aggcttcttc acctcagccc cagactgcac cagctgcacc tg 342

<210> 54
<211> 366
<212> DNA
<213> Homo sapiens

<400> 54
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tcctgtgcag cctctggatt cacctttagc aactttgcca tgagctgggt ccgccaggct 120

cccgggaagg ggctggagtg ggtcgcggct attggcggta gaagtggtac cacattctac 180

gcggactccg tgaagggccg gttcaccatc tccagagaca attccaagaa cacggtctat 240

ctggaaatga acagtctgag agccgaggac acagccattt attactgtgc gaaaagaggg 300

cgcgggggt ataagtatta tgggatggac gtctggggcc aggggaccac ggtcaccgtg 360

tcgagt 366

<210> 55
<211> 366
<212> DNA
<213> Homo sapiens

<400> 55
actcgacacg gtgaccgtgg tcccctggcc ccagacgtcc atcccataat acttataccc 60

cccgcgccct cttttcgcac agtaataaat ggctgtgtcc tcggctctca gactgttcat 120

ttccagatag accgtgttct tggaattgtc tctggagatg gtgaaccggc ccttcacgga 180

gtccgcgtag aatgtggtac cacttctacc gccaatagcc gcgacccact ccagcccctt 240

cccgggagcc tggcggaccc agctcatggc aaagttgcta aaggtgaatc cagaggctgc 300

acaggagagt ctcagggacc ccccaggctg taccaagtct cccccagact ccaccagctg 360

cacctc 366